

IN THE CLAIMS:

Cancel claim 2.

1. (currently amended) A purified polypeptide comprising an amino acid sequence selected from the group consisting of:
 - a) ~~the~~ an amino acid sequence as set forth in ~~of~~ SEQ ID NO:2,and
 - b) an amino acid sequence ~~a fragment of SEQ ID NO:1~~ comprising ~~the kinesin motor domain from~~ amino acid residue 1 to amino acid residue 340 of SEQ ID NO:2.
2. (cancelled)
3. (currently amended) A pharmaceutical composition comprising a the polypeptide of claim 1 and a pharmaceutically acceptable excipient.
4. (original) A composition of claim 3, wherein the polypeptide has the sequence of SEQ ID NO:2.
5. (currently amended) A method for screening a compound for effectiveness as an agonist of a the polypeptide of claim 1, ~~the method~~ comprising:
 - a) exposing a sample comprising a the polypeptide of claim 1 to a compound, and
 - b) detecting agonist activity in the sample.
6. (currently amended) A method for screening a compound for effectiveness as an antagonist of a the polypeptide of claim 1, the method comprising:
 - a) exposing a sample comprising a polypeptide of claim 1 to a compound, and
 - b) detecting antagonist activity in the sample.

7. (currently amended) An isolated ~~and purified~~ polynucleotide nucleic acid molecule comprising a sequence of nucleotides encoding a polypeptide comprising an amino acid sequence as set forth in of SEQ ID NO:1.

8. (currently amended) An isolated ~~and purified~~ polynucleotide which hybridizes under conditions of 250 mM NaCl, 25 mM trisodium citrate, 1% SDS, 50% formamide and 200 µg/ml ssDNA at 42°C., and wash conditions of 15 mM NaCl, 1.5 mM trisodium citrate, and 0.1% SDS at 68°C to the sequence of nucleotides as set forth in ~~polynucleotide of claim 7.~~

9. (currently amended) A method for detecting a nucleic acid molecule having a sequence of nucleotides substantially similar to the nucleic acid molecule of claim 7 ~~polynucleotide~~, the method comprising the steps of:

a) hybridizing the nucleic acid molecule ~~polynucleotide~~ of claim 7 to at least one nucleic acid in a sample, under conditions favoring the formation of ~~thereby forming~~ a hybridization complex; and

b) detecting the hybridization complex, wherein said hybridization is performed at 42°C in a solution containing 250 mM NaCl, 25 mM trisodium citrate, 1% SDS, 50% formamide and 200 µg/ml ssDNA followed by washing at 68°C in a solution of 15 mM NaCl, 1.5 mM trisodium citrate, and 0.1% SDS wherein the presence of the hybridization complex correlates with the presence of the polynucleotide in the sample.

10. (original) The method of claim 9 further comprising amplifying the polynucleotide prior to hybridization.

11. (currently amended) An isolated ~~and purified~~ polynucleotide nucleic acid molecule comprising a sequence of nucleotides that encode a polypeptide comprising the amino acid sequence as set forth in SEQ ID NO: 2 ~~comprising the polynucleotide sequence of SEQ ID NO:1.~~

12. (currently amended) An expression vector comprising the ~~polynucleotide~~ nucleic acid molecule of claim 7.

12. 13. (original) A host cell comprising the expression vector of claim

14. (currently amended) A method for producing a the polypeptide of claim 1, ~~the method~~ comprising the steps of:

- a) culturing the host cell of claim 13 under conditions suitable for the expression of the polypeptide; and
- b) recovering the polypeptide from the host cell culture.

15. (currently amended) A method ~~Method~~ of modulating cellular proliferation in a mammal in need thereof comprising administering to said mammal an amount of the a pharmaceutical composition of claims 3 effective to modulate cellular proliferation, said composition comprising a pharmaceutically acceptable vehicle and a HsCENP-E protein characterized as having an ATP binding site, and a motor domain comprising and an amino acid sequence from amino acid at position 1 through amino acid at position 340 as set forth in SEQ ID NO:2.

16. (currently amended) A method for inhibiting HsCENP-E mediated/induced cellular proliferation of a cell in culture, said method comprising the steps of:

- a) providing an oligonucleotide comprising at least 18 contiguous nucleotide bases which are ~~perfectly~~ complementary to a nucleotide base sequence region contained in a nucleic acid sequence as set forth in SEQ ID NO.1, and
- b) contacting said cell with said oligonucleotide under conditions such that said oligonucleotide is delivered within said cell and hybridizes with said nucleotide base sequence region, thereby inhibiting HsCENP-E mediated/induced cellular proliferation of said cell.

17. (currently amended) A method of detecting the presence of cancer in an individual comprising:

- (a) obtaining a biological sample from said individual;

- (b) incubating said biological sample with at least one antibody which is immunoreactive with a gene product encoded by the nucleic acid molecule of Claim 7, ~~comprising the nucleotide sequence as set forth in SEQ ID NO:1~~
- (c) detecting immunoconjugates which form as a consequence of the incubation of step (b); and
- (d) relating the amount of immunoconjugates of step (c) to the presence of cancer, wherein cancer is present when said amount is greater than a threshold value.

18. (currently amended) ~~An isolated and~~ The substantially purified polypeptide of claim 1, wherein said polypeptide comprises an amino acid sequence as set forth in SEQ ID NO:2 ~~encoded by the nucleotide sequence of SEQ ID NO:1.~~

Please add new claims as follows:

19 (New) The substantially purified polypeptide of claim 1, wherein said polypeptide comprises an amino acid sequence comprising amino acids at position 1 through 340 of SEQ ID NO:2 .